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FILE 'HOME' ENTERED AT 17:00:54 ON 03 MAR 2004

=> file agricola biosis embase caplus

COST IN U.S. DOLLARS	SINCE FILE ENTRY	TOTAL SESSION
FULL ESTIMATED COST	0.21	0.21

FILE 'AGRICOLA' ENTERED AT 17:01:06 ON 03 MAR 2004

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=> s hemoglobin and seeds? and breeding
L1 2 HEMOGLOBIN AND SEEDS? AND BREEDING

=> d 11 1-2

L1 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2004 ACS on STN
AN 2000:15341 CAPLUS
DN 132:60124
TI Nonsymbiotic plant ***hemoglobins*** to maintain cell energy status in transgenic systems
IN Guy, Phillip; Duff, Stephen; Xianzhou, Nie; Hill, Robert; Durnin, Douglas; Sowa, Aleksander
PA University of Manitoba, Can.
SO PCT Int. Appl., 44 pp.
CODEN: PIXXD2
DT Patent
LA English
FAN.CNT 1

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2000000597	A2	20000106	WO 1999-CA587	19990624
	WO 2000000597	A3	20000323		

W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS,

=> s hemoglobin and seeds? and selection

L2 1 HEMOGLOBIN AND SEEDS? AND SELECTION

=> d 12 1

L2 ANSWER 1 OF 1 EMBASE COPYRIGHT 2004 ELSEVIER INC. ALL RIGHTS RESERVED.
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AN 94324312 EMBASE

DN 1994324312

TI Trace element nutrition during pregnancy.

AU Wada L.; King J.C.

CS Department of Nutritional Sciences, College of Natural Resources,
University of California, Berkeley, CA 94720, United States

SO Clinical Obstetrics and Gynecology, (1994) 37/3 (574-586).

ISSN: 0009-9201 CODEN: COGYAK

CY United States

DT Journal; Conference Article

FS 010 Obstetrics and Gynecology

017 Public Health, Social Medicine and Epidemiology

LA English

SL English

=> d 12 1 ab

L2 ANSWER 1 OF 1 EMBASE COPYRIGHT 2004 ELSEVIER INC. ALL RIGHTS RESERVED.
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AB The best means to ensure an optimal intake of trace elements during
pregnancy is the consumption of a well-balanced diet that includes both
animal- and plant-food sources. Although vegetarian diets provide
reasonable sources of trace elements, especially in dried beans and
seeds, flesh foods contain higher concentrations of trace
elements

that are in a more readily absorbable form. Iron is the only trace element
for which routine supplementation is recommended. In the United States,
the iodine content of the food supply is sufficiently high to make
supplementation unnecessary, but use of iodized salt is not
contraindicated. There is no need to screen pregnant women routinely for
trace element status, except to monitor ***hemoglobin*** and
hematocrit as an indicator of iron status. The best indicators that a
woman's trace element status may be at risk is a history of poor food
selections, a clinical disorder that alters trace element use or
excretion, or prior residence in a region of the world where the trace
element content of the soil is low.

=> s hemoglobin and seed? and selection

L3 10 HEMOGLOBIN AND SEED? AND SELECTION

=> duplicate remove l3

DUPLICATE PREFERENCE IS 'AGRICOLA, BIOSIS, EMBASE, CAPLUS'

KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n

PROCESSING COMPLETED FOR L3

L4 8 DUPLICATE REMOVE L3 (2 DUPLICATES REMOVED)

While the ***seed*** and the pollen carry species characters, they also transmit certain obscure properties which are peculiar to each of the sex elements; this was shown by differences in the offspring of reciprocal crosses. The germplasm is recognized as a stereochem. system. Such phenomena as sports, reversion, variation, and the genesis of strains, subspecies, and species are due to gradual and progressive changes and ultimate fixation; hybridization is probably the chief method in the genesis of species.

=> s selection and seed and nonsymbiotic and hemoglobin
L5 0 SELECTION AND SEED AND NONSYMBIOTIC AND HEMOGLOBIN

=>
=> seed(w)selection and breeding
SEED(W) SELECTION IS NOT A RECOGNIZED COMMAND
The previous command name entered was not recognized by the system.
For a list of commands available to you in the current file, enter
"HELP COMMANDS" at an arrow prompt (=>).

=> s seed(w) selection and breeding
L6 52 SEED(W) SELECTION AND BREEDING

=> duplicate remove 16
DUPLICATE PREFERENCE IS 'AGRICOLA, BIOSIS, EMBASE, CAPLUS'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/ (N) :n
PROCESSING COMPLETED FOR L6
L7 49 DUPLICATE REMOVE L6 (3 DUPLICATES REMOVED)

=> d 17 1-10 ti

L7 ANSWER 1 OF 49 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
TI Selection for large seed and high protein in two- and three-parent soybean populations.

L7 ANSWER 2 OF 49 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
DUPLICATE 1
TI Genetic effects on grain shape traits of indica black pericarp rice and their genetic correlations with main mineral element contents in grains.

L7 ANSWER 3 OF 49 CAPLUS COPYRIGHT 2004 ACS on STN
TI System for functional gene discovery in plants using multigeneration ***breeding*** with database tracking of phenotypes

L7 ANSWER 4 OF 49 CAPLUS COPYRIGHT 2004 ACS on STN
TI Non-random selection of livestock and crops and tissue products in ***breeding*** for uniformity using phenomics and bioinformatics

L7 ANSWER 5 OF 49 CAPLUS COPYRIGHT 2004 ACS on STN
TI Selection for high seed oil content in soybean families derived from plants regenerated from protoplasts and tissue cultures

L7 ANSWER 6 OF 49 CAPLUS COPYRIGHT 2004 ACS on STN
TI Selection of soybean mutants with increased concentrations of seed methionine and cysteine

L7 ANSWER 7 OF 49 CAPLUS COPYRIGHT 2004 ACS on STN

cultivar Idared, through parthenogenesis in situ.
AU De Witte, Krista; Keulemans, Johan
CS Fruitteeltcentrum K.U. Leuven, Willem de Croylaan 42, B-30001
Leuven-Heverlee, Belgium
SO Euphytica, (1994) Vol. 77, No. 1-2, pp. 141-146.
CODEN: EUPHAA. ISSN: 0014-2336.
DT Article
LA English
ED Entered STN: 25 Jan 1995
Last Updated on STN: 25 Jan 1995

=> FIL STNGUIDE	SINCE FILE	TOTAL
COST IN U.S. DOLLARS	ENTRY	SESSION
FULL ESTIMATED COST	93.91	94.12
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)	SINCE FILE	TOTAL
CA SUBSCRIBER PRICE	ENTRY	SESSION
	-2.08	-2.08

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AND TECHNOLOGY CORPORATION, AND FACHINFORMATIONSZENTRUM KARLSRUHE

FILE CONTAINS CURRENT INFORMATION.
LAST RELOADED: Feb 27, 2004 (20040227/UP).

=> s single(w) seed(w) descent
221 SINGLE
0 SEED
0 DESCENT
L8 0 SINGLE(W) SEED(W) DESCENT

=> file agricola biosis embase caplus	SINCE FILE	TOTAL
COST IN U.S. DOLLARS	ENTRY	SESSION
FULL ESTIMATED COST	0.12	94.24
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)	SINCE FILE	TOTAL
CA SUBSCRIBER PRICE	ENTRY	SESSION
	0.00	-2.08

FILE 'AGRICOLA' ENTERED AT 17:31:22 ON 03 MAR 2004

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L10 ANSWER 3 OF 341 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
AB, This paper reviews some measurements of representativeness such as the effective population size (N_e) useful in genetic resources conservation and plant breeding research. Some easy-to-use expressions for estimating N_e in a number of practical conditions are given. For the case of several subpopulations, N_e is scarcely influenced by the total inbreeding. Thus, N_e becomes mostly dependent on the allelic diversity among subpopulations and the number of subpopulations sampled. When, under natural conditions, levels of interpopulation allelic divergence are low, it is misleading to believe that a small number of subpopulations will be sufficient to attain adequate effective population sizes. When a single population is considered and seeds being sampled have a family structure, at the limit N_e depends only on the number of seed parents and the coancestry coefficient among sibs within families. Accession regeneration is the case where the reference population is of finite size. Gametic control is a major factor in regeneration. The loss of up to 20% of seeds may be recovered in terms of N_e when female gametic control is applied. This is not attainable with random sampling of seeds. When studying N_e in recurrent selection schemes, results showed that the gain in N_e through gametic control is very small when selection is intensive. When comparing effective population sizes for the ***single*** ***seed***
descent (SSD) method versus the bulk system, results showed that SSD maintains genetic drift at a low level and offers a much better protection against random loss of alleles during selfing generations. Estimating population parameters, through codominant genetic markers is fundamental for obtaining reliable estimates of effective population size.

=> s single(w) seed(w) descent and selection and trait and breeding
L11 52 SINGLE(W) SEED(W) DESCENT AND SELECTION AND TRAIT AND BREEDING

=> duplicate remove l11
DUPLICATE PREFERENCE IS 'AGRICOLA, BIOSIS, EMBASE, CAPLUS'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n

PROCESSING COMPLETED FOR L11
L12 38 DUPLICATE REMOVE L11 (14 DUPLICATES REMOVED)

=> d l12 1-15

L12 ANSWER 1 OF 38 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
DUPLICATE 1
AN 2003:578040 BIOSIS
DN PREV200300582830
TI Mapping QTL for ***traits*** associated with resistance to ferrous iron toxicity in rice (*Oryza sativa* L.), using japonica chromosome segment substitution lines.
AU Wan Jian-Lin [Reprint Author]; Zhai Hu-Qu; Wan Jian-Min [Reprint Author]; Yasui, Hideshi; Yoshimura, Atsushi
CS State Key laboratory of Crop Genetics and Germplasm Enhancement, Jiangsu Plant Gene Engineering Research Center, Nanjing Agricultural University, Nanjing, 210095, China
wanjm@mail.njau.edu.cn
SO Acta Genetica Sinica, (Oct 2003) Vol. 30, No. 10, pp. 893-898. print.
ISSN: 0379-4172 (ISSN print).
DT Article

SSD, 34.5% of the original population was lost due to barrenness at the high density in 1 cross and 26.5% in the other. Seed of surviving genotypes was increased for a replicated field trial where maturity (days after 1 June), plant height (cm) oil content (NMR value), and grain yield (g/hill plot) were measured. Means and variances between progenies advanced under high or low densities generally did not differ for any of the ***traits***, leading to the conclusion that the genotypes eliminated due to barren plants were randomly and not selectively eliminated due to ***selection*** pressures of the technique.

L12 ANSWER 38 OF 38 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
AB Heredity of a few ***traits*** relevant to plant development and assimilation area was studied in Toulouse, France, in 2 independent non-selected hybrid populations segregating in F3 and F4. Both generations were grown related to each other, as well as to the initial F2, by ***single*** ***seed*** ***descent***. The F4 material was also analyzed on a plant mean basis in 2 replicated rows at normal planting density. For both populations as isolated plants, heritability values were consistently high (0.32-0.70) for leaf area and earliness. They were low (< 1) for 2nd development phase duration and weight per seed. The 2 populations did not differ much from each other, except for the heritability size of the various characters. When the F4 was grown under conditions of plant competition, the heritability values remained roughly the same, but for the number of pods per plant it slightly decreased. The expected genetic progress resulting, at normal density, from a ***selection*** performed among spaced plants, showed more spread differences between characters, with total similarity between both families. The more responsive ***trait*** was leaf area, followed by yield. Although the 2 characters are positively correlated, it is possible to associate, in a ***breeding*** program, an improved yield with a moderate leaf area, that is recommendable in the climatic conditions of France.

=> s l12 and protein
L13 4 L12 AND PROTEIN.

=> d l13 1-4 ibib ab

L13 ANSWER 1 OF 4 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.
(2004) on STN

ACCESSION NUMBER: 2000:31462 AGRICOLA
DOCUMENT NUMBER: IND22034394
TITLE: Agronomic performance of lines derived from anther culture, maize pollination and ***single*** - ***seed*** ***descent*** in a spring wheat cross.
AUTHOR(S): Ma, H.; Busch, R.H.; Riera-Lizarazu, O.; Rines, H.W.; Dill-Macky, R.
CORPORATE SOURCE: University of Minnesota, St. Paul, MN.
AVAILABILITY: DNAL (442.8 Z8)
SOURCE: Theoretical and applied genetics, Aug 1999. Vol. 99, No. 3/4. p. 432-436
Publisher: Berlin; Springer-Verlag
CODEN: THAGA6; ISSN: 0040-5752

Concomitant with our QTL identification and manipulation efforts are several ongoing expts. based on functional genomics. We are convinced that emerging technologies based on gene expression have the potential to provide complementary information that will promote more efficient germplasm improvement. Indeed, understanding the genetic basis of the essential physiol. parameters of drought tolerance in maize, together with the data generated by profiling expts., should allow the identification of the key pathways involved in drought stress and further our understanding of how they interact. This, in turn, will lead to develop strategies to improve the tolerance of maize and other crops to water-limited conditions.

REFERENCE COUNT: 38 THERE ARE 38 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> protein(w) content and seed and selection and breeding and plant
PROTEIN(W) CONTENT IS NOT A RECOGNIZED COMMAND
The previous command name entered was not recognized by the system.
For a list of commands available to you in the current file, enter
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=> s protein(w) content and seed and selection and breeding and plant
L14 281 PROTEIN(W) CONTENT AND SEED AND SELECTION AND BREEDING AND PLANT

=> s l14 and barley
L15 17 L14 AND BARLEY

=> duplicate remove l15
DUPLICATE PREFERENCE IS 'BIOSIS, CAPLUS'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/ (N) :n
PROCESSING COMPLETED FOR L15
L16 17 DUPLICATE REMOVE L15 (0 DUPLICATES REMOVED)

=> d l16 1-10 ti

L16 ANSWER 1 OF 17 CAPLUS COPYRIGHT 2004 ACS on STN
TI High protein phenotype-associated ***plant*** genes and their use for generating transgenic ***plants*** with improved nutritional properties

L16 ANSWER 2 OF 17 CAPLUS COPYRIGHT 2004 ACS on STN
TI ***Barley*** malting quality and yield interrelationships and the effect on yield distribution of ***selection*** for malting quality in the early generations

L16 ANSWER 3 OF 17 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
TI A synthetic hexaploid ($2n = 42$) oat from the cross of *Avena strigosa* ($2n = 14$) and domesticated *A. magna* ($2n = 28$).

L16 ANSWER 4 OF 17 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
TI ***Protein*** ***content*** heritability in ***selection*** of ***barley*** based on intraplant variation in kernel anatomical structure.

L16 ANSWER 5 OF 17 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
TI ***Selection*** of Dry pea (*Pisum sativum*) varieties and their performance by comparison with ***barley*** and wheat in Mediterranean

protein ***content*** . However, some differences in
protein ***content*** were significant, e.g. grains 11, 15,
and 17 contained 12, 13.8, and 15.7% protein, resp. Grains of the KM-1192
cultivar and its F1 and F2 hybrids pollinated by Hyproly contained
8.7-12.4, 9.3-11.0, and 6.9-12.4% protein, resp. Thus, the variability
increased in F2, whereas in F1 it resembled that of the mother cultivar.

=> single(w)trait(w)variability and protein and seed and barley and selection
SINGLE(W) TRAIT(W) VARIABILITY IS NOT A RECOGNIZED COMMAND

The previous command name entered was not recognized by the system.

For a list of commands available to you in the current file, enter
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=> s single(w)trait(w)variability and protein and seed and barley and selection
L17 0 SINGLE(W) TRAIT(W) VARIABILITY AND PROTEIN AND SEED AND BARLEY
AND SELECTION

=> s single(w)trait(w)variability and protein and seed and selection
L18 0 SINGLE(W) TRAIT(W) VARIABILITY AND PROTEIN AND SEED AND SELECTION
N

=> s single(w)trait(w)variability and seed and selection
L19 0 SINGLE(W) TRAIT(W) VARIABILITY AND SEED AND SELECTION

=> s single(w)trait(w)variability and seed and protein and selection
L20 0 SINGLE(W) TRAIT(W) VARIABILITY AND SEED AND PROTEIN AND SELECTION
N

=> s single(w)trait and seed and protein and selection
L21 1 SINGLE(W) TRAIT AND SEED AND PROTEIN AND SELECTION

=> d l21 ibib ab

L21 ANSWER 1 OF 1 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
ACCESSION NUMBER: 1997:242957 BIOSIS

DOCUMENT NUMBER: PREV199799542160

TITLE: ***Selection*** for yield, ***protein*** , and oil
in soybean crosses between adapted and introduced parents.

AUTHOR(S): Scott, Roy A. [Reprint author]; Kephart, Kevin D.

CORPORATE SOURCE: Plant Sci. Dep., South Dakota State Univ., Box 2140C, NPB
247, Brookings, SD 57007, USA

SOURCE: Field Crops Research, (1997) Vol. 49, No. 2-3, pp. 177-185.
ISSN: 0378-4290.

DOCUMENT TYPE: Article

LANGUAGE: English

ENTRY DATE: Entered STN: 13 Jun 1997

Last Updated on STN: 13 Jun 1997

AB Soybean (*Glycine max* (L.) Merr.) breeders can sometimes increase genetic
variability in their germplasm through wide crosses with plant
introduction (PI) material, and in that way lead to slow progress in
developing high-yielding cultivars. A study was conducted to evaluate the
potential of crosses of adapted times introduced soybean lines for
improving yield and ***protein*** and oil concentrations. Five
soybean cultivars adapted to South Dakota were crossed in selected
combinations with four PI lines to develop F₃-derived lines. The adapted
and PI lines were selected for wide variation in ***protein*** and oil